SEQUENCE LISTING

<110> Ferrick, David A.
      Swift, Susan E.
      Armstrong, Randall
      Fox, Bryan

<120> Methods and Compositions for Screening for Modulators and Ige Synthesis, Secretion and Switch Rearrangement

<130> A-66038-4/RMS/JJD/DLR

<140> US 09/966,976
<141> 2001-09-27

<150> US 09/076,624
<151> 1998-05-12

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tcatgccatc cgtaagatgc ttttctgtga ctggtgagta ctcaaccaag tcattctgag    7920

aatagtgtat gcggcgaccg agttgctctt gcccggcgtc aacacgggat aataccgcgc    7980

cacatagcag aactttaaaa gtgctcatca ttggaaaacg ttcttcgggg cgaaaactct    8040

caaggatctt accgctgttg agatccagtt cgatgtaacc cactcgtgca cccaactgat    8100

cttcagcatc ttttactttc accagcgttt ctgggtgagc aaaaacagga aggcaaaatg    8160

ccgcaaaaaa gggaataagg gcgacacgga aatgttgaat actcatactc ttcctttttc    8220

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gt    8282
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<210> 8
<211> 8345
<212> DNA
<213> Artificial sequence

17

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ttacggggtc attagttcat agcccatata tggagttccg cgttacataa cttacggtaa   180

atggcccgcc tggctgaccg cccaacgacc cccgcccatt gacgtcaata atgacgtatg   240

ttcccatagt aacgccaata gggactttcc attgacgtca atgggtggag tatttacggt   300

aaactgccca cttggcagta catcaagtgt atcatatgcc aagtacgccc cctattgacg   360

tcaatgacgg taaatggccc gcctggcatt atgcccagta catgacctta tgggactttc   420

ctacttggca gtacatctac gtattagtca tcgctattac catggtgatg cggttttggc   480

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gcagagctca ataaaagagc ccacaacccc tcactcgggg cgccagtcct ccgattgact   720

gagtcgcccg ggtacccgtg tatccaataa accctcttgc agttgcatcc gacttgtggt   780

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gctctgcaga atggccaacc tttaacgtcg gatggccgcg agacggcacc tttaaccgag  1500

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18

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agactaagaa cctagaacct cgctggaaag gaccttacac agtcctgctg accaccccca    2040

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20

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21

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22

<210> 9
<211> 61
<212> PRT
<213> Artificial sequence

<220>
<223> coiled-coil presentation structure

<400> 9

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Met Gly Cys Ala Ala Leu Glu Ser Glu Val Ser Ala Leu Glu Ser Glu
1               5                   10                  15


Val Ala Ser Leu Glu Ser Glu Val Ala Ala Leu Gly Arg Gly Asp Met
            20                  25                  30


Pro Leu Ala Ala Val Lys Ser Lys Leu Ser Ala Val Lys Ser Lys Leu
        35                  40                  45


Ala Ser Val Lys Ser Lys Leu Ala Ala Cys Gly Pro Pro
    50                  55                  60
```

<210> 10
<211> 6
<212> PRT
<213> Artificial sequence

<220>
<223> looped structure of coiled-coil presentation structure

<400> 10

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Gly Arg Gly Asp Met Pro
1               5
```

<210> 11
<211> 69
<212> PRT
<213> Artificial sequence

<220>
<223> minibody presentation structure

<400> 11

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Met Gly Arg Asn Ser Gln Ala Thr Ser Gly Phe Thr Phe Ser His Phe
1               5                   10                  15


Tyr Met Glu Trp Val Arg Gly Gly Glu Tyr Ile Ala Ala Ser Arg His
            20                  25                  30
```

Lys His Asn Lys Tyr Thr Thr Glu Tyr Ser Ala Ser Val Lys Gly Arg
         35   40   45

Tyr Ile Val Ser Arg Asp Thr Ser Gln Ser Ile Leu Tyr Leu Gln Lys
         50   55   60

Lys Lys Gly Pro Pro
65

<210> 12
<211> 7
<212> PRT
<213> Simian virus 40

<400> 12

Pro Lys Lys Lys Arg Lys Val
1   5

<210> 13
<211> 6
<212> PRT
<213> Homo sapiens

<400> 13

Ala Arg Arg Arg Arg Pro
1   5

<210> 14
<211> 10
<212> PRT
<213> Mus musculus

<400> 14

Glu Glu Val Gln Arg Lys Arg Gln Lys Leu
1   5   10

<210> 15
<211> 9
<212> PRT
<213> Mus musculus

<400> 15

Glu Glu Lys Arg Lys Arg Thr Tyr Glu
1   5

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<210> 16
<211> 20
<212> PRT
<213> Xenopus laevis

<400> 16

Ala Val Lys Arg Pro Ala Ala Thr Lys Lys Ala Gly Gln Ala Lys Lys
1               5                   10                  15


Lys Lys Leu Asp
            20


<210> 17
<211> 10
<212> PRT
<213> Artificial sequence

<220>
<223> stability sequence

<220>
<221> MISC_FEATURE
<222> (3)..(6)
<223> "Xaa" at positions 3 to 6 can be any amino acid.


<400> 17

Met Gly Xaa Xaa Xaa Xaa Gly Gly Pro Pro
1               5                   10


<210> 18
<211> 5
<212> PRT
<213> Artificial sequence

<220>
<223> linker sequence

<400> 18

Gly Ser Gly Gly Ser
1               5


<210> 19
<211> 4
<212> PRT
<213> Artificial sequence

<220>
<223> linker sequence
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<400> 19

Gly Gly Gly Ser
1